

6



OIEP

RAW SEQUENCE LISTING

DATE: 04/08/2002

PATENT APPLICATION: US/10/015,219

TIME: 08:55:08

Input Set : A:\493c1.app.txt

Output Set: N:\CRF3\04082002\J015219.raw

P.S

ENTERED

4 <110> APPLICANT: Algate, Paul A.
 6 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
 7 AND DIAGNOSIS OF OVARIAN CANCER
 10 <130> FILE REFERENCE: 210121.493C1
 12 <140> CURRENT APPLICATION NUMBER: US 10/015,219
 13 <141> CURRENT FILING DATE: 2002-03-02
 15 <160> NUMBER OF SEQ ID NOS: 1739
 17 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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 20 <211> LENGTH: 264
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Homo sapiens
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 26 ctgatatccg ggactttttc cctttcacct tccattatag gatgatgcag caagaaggcc 120
 27 ctaccagat accagtcctt tgatcttgga ctaccaggc ttcagaacta taagaaataa 180
 28 atctccgttc tttgaggatt acccagtttg tgctattctg ttatggcagc acaaaatgga 240
 29 ctaagacata ggtccaatct aagg 264
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 34 <213> ORGANISM: Homo sapiens
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 37 <221> NAME/KEY: misc_feature
 38 <222> LOCATION: 9, 10, 494, 534
 39 <223> OTHER INFORMATION: n = A,T,C or G
 41 <400> SEQUENCE: 2
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 43 cattttgagc aatggggaac gctcacggac tgtgtggtta tgagagatcc aaacaccaag 120
 44 cgctccaggg gctttgggtt tgcacatat gccactgttg aggaggtgga tgcagctatg 180
 45 aatgcaaggc cacacaaggt ggatggaaga gttgtggaac caaagagagc tgtctccaga 240
 46 gaagattctc aaagaccagg tgcccactta actgtgaaaa agatatttgt tgggtggcatt 300
 47 aaagaagaca ctgaagaaca tcacctaaaga gattattttg aacagtatgg aaaaattgaa 360
 48 gtgattgaaa tcatgactga ccgaagcagt ggcaagaaaa ggggctttgc ctttgtaacc 420
 49 tttgacgacc atgactccgt ggataagatt gtcattcaga aatccattcc tgtgaatgga 480
 50 cctgccccgc cggncaaagg cgaaattcaa cacactttgg cggcggttacc taanggatcc 540
 51 caacttcggt 550
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 54 <211> LENGTH: 434
 55 <212> TYPE: DNA
 56 <213> ORGANISM: Homo sapiens
 58 <400> SEQUENCE: 3
 59 aaatttatga aactttcgaa cagtagcaac tgaaatttgt cacttttctg ttacgcagag 60

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61 ttgtcagcac tacatacatc ttttttttgc ggggggcggg ggggacagag tctcactgtg 180
62 tcaactcagac tggagtacag tgatgcgatac tcggctcact gcaacctccg cctcctgggt 240
63 tcaagcgatt ctcctgcttc agcctcctga gtagctggga ttacaggtgc acaccaccac 300
64 gccccgctaa tttttgtatt tttagtagag atgggggttc accatgttgg tcaggctggg 360
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66 accaccgcac ctgg                                     434
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69 <211> LENGTH: 381
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75 gccccaaatg ccaggctgca ctgatattta ttggatataa gacaaagggg cagggttaagg 120
76 aatgtgaacc atctccaata ataggtaagg tcacatgggt catgtgtcca ctggacaggg 180
77 ggcccttccc tgccctggcag cagaggcaga gagagagaga agagagagag acagcttatg 240
78 ccattatttc tgcataatcag acatttagta ctttcactaa tttgctcctg ctatctaaaa 300
79 ggcagagcca ggtatacagg atggaacatg aaagcggact aggagcgtga ccactgaagc 360
80 acagcatcac agggagacag g                                     381
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83 <211> LENGTH: 422
84 <212> TYPE: DNA
85 <213> ORGANISM: Homo sapiens
87 <400> SEQUENCE: 5
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89 ggcctgccat gcttaaaata ttaactctct ggccctttta gaaaaaaacg tgctgacccc 120
90 tgctctagat caaagaaaac aaacctcaaa aatactttcc tccctctacc ccacttgacc 180
91 cttgtcccgg ggcagtaggc atctccgtca aaactcttgt ccttggtctg tggtaacttt 240
92 ctcagctccc caacccatgt cctcaaaagt cccctcccta tagggcaaga acccagcaac 300
93 ttcgctctgc cccgactcta ggcgggatgt agctcatttt gggatacagag tctccatcgt 360
94 ggagcctggc ttcttccgaa cccctgtgac caacctggag agtctggaga aaacctgca 420
95 gg                                     422
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98 <211> LENGTH: 261
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103 <221> NAME/KEY: misc_feature
104 <222> LOCATION: 148, 180
105 <223> OTHER INFORMATION: n = A,T,C or G
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109 tctgtcttgt tgcaaagcca gcacagcaag cagcctccgc attagttcca tagcttgact 120
110 ggcttctaag atgggcatg tcaagatnca gaaatctcaa agcatccctt ctttgggctn 180
111 catcatccaa gggtagaaaa cagcagagcc taagttagaa gtctgagtca acaccttggc 240
112 tcagttttca aaatgaattt t                                     261
114 <210> SEQ ID NO: 7
115 <211> LENGTH: 428
116 <212> TYPE: DNA

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117 <213> ORGANISM: Homo sapiens
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121 ctaaaaaacg caacaatcag atacggaaga cctcttatgc tcagcaccaa caggtccgcc 120
122 aaatccggaa gaaagatgat ggaaatcatg acccgagagg tgcagacaaa tgacttgaaa 180
123 gaagtgggtca ataaattgat tccagacagc attggaaaag acatagaaaa ggcttgccaa 240
124 tctattttatc ctctccatga tgtcttcgtt agaaaagtaa aaatgctgaa gaagcccaag 300
125 tttgaattgg gaaagctcat ggagcttcat ggtgaaggca gtagttctgg aaaagccact 360
126 ggggacgaga caggtgctaa agttgaacga gctgatggat atgaaccacc agtccaagaa 420
127 atctgttt 428
129 <210> SEQ ID NO: 8
130 <211> LENGTH: 305
131 <212> TYPE: DNA
132 <213> ORGANISM: Homo sapiens
134 <400> SEQUENCE: 8
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136 gttgctgtct cgcttggag aaatcgtga ggaaggctca gagactcagg tcccaggact 120
137 ggacatatct tgcactcctg ccctctgact tcagccgcta cttccaatat gaggggtctc 180
138 tgactacacc gccctgtgcc caggggtgca tctggactgt gtttaaccag acagtgatgc 240
139 tgagtgctaa gcagctccac accctctctg acaccctgtg gggacctggg gactctcggc 300
140 tacag 305
142 <210> SEQ ID NO: 9
143 <211> LENGTH: 344
144 <212> TYPE: DNA
145 <213> ORGANISM: Homo sapiens
147 <400> SEQUENCE: 9
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149 gaaacacttt cctacagacc catccaaatg tgtggagtac aatgcactgt gagatctgtg 120
150 tatggtgtgt taataacaat aagaaactta gggaagcagg ctgtggactt ctggaattac 180
151 caacaggaat gaggaagaa gaaaactgga gtttccagtc tctgagttct acctgatgta 240
152 actcttgatt ggttttaaga actttgttgg ccttcatttc atatctgact gcaagctgat 300
153 ttttctttct tgctttcatt ttaattagtc caaaattaag tttt 344
155 <210> SEQ ID NO: 10
156 <211> LENGTH: 377
157 <212> TYPE: DNA
158 <213> ORGANISM: Homo sapiens
160 <400> SEQUENCE: 10
161 aaaaccttta gcatttctgc ctataatatt tgggttttct tcttttctta tctttatttg 60
162 ataagtccca tcaaatattt tcccataat cacaatgttt tcttttctact ttgctcaaga 120
163 actgagttat gagctccaaa atttgacaa actctacatt ggctaagttt tagtcatttg 180
164 cactgctaag aaagatgaca attcagcatg ctgaagatga cttcctccct tataaagggg 240
165 ctaacacaga gggcaatact gttcatgctt ctgattcttg atcacaagaa ttgctttagg 300
166 caattacaat catgtctcct ctgacacatc atattattca agtgagacag agaaagaaga 360
167 tgtcctatgt cacacag 377
169 <210> SEQ ID NO: 11
170 <211> LENGTH: 381
171 <212> TYPE: DNA
172 <213> ORGANISM: Homo sapiens
174 <400> SEQUENCE: 11

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176 gcccacaaatg ccaggctgca ctgatattta ttggatataa gacaaagggg cagggtaagg 120
177 aatgtgaacc atctccaata ataggtaagg tcacatgggt catgtgtcca ctggacaggg 180
178 ggcccttccc tgccctggcag cagaggcaga gagagagaga agagagagag acagcttatg 240
179 ccattatttc tgcatatcag acatttagta ctttactaa tttgtcctg ctatctaaaa 300
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181 acagcatcac agggagacag g                                     381
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186 <213> ORGANISM: Homo sapiens
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190 gggccaaagt ggaggccggg tcagtgtgga ggtggattcc gctccgggca ccgatctcgc 120
191 caagatcctg agtggcatgc gaagccaata tgaggatcat gccgagcaga accggaagga 180
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195 <211> LENGTH: 355
196 <212> TYPE: DNA
197 <213> ORGANISM: Homo sapiens
199 <220> FEATURE:
200 <221> NAME/KEY: misc_feature
201 <222> LOCATION: 344, 348
202 <223> OTHER INFORMATION: n = A,T,C or G
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206 tgattattta catttttagta attggacaat cccggctcag gaggagggtg caagaatctg 120
207 caaaagttgg agggagcggc ccaggagaac aaacagcaag ccttatttcc cctagcccat 180
208 cccccaaaaa accatccatc ccattcctagt gtctgggtgt gtcgggtgtg gtccatcttc 240
209 cattccttcc caaattatgg aagtaagggt cttctcacca gaataagagc acttgggata 300
210 acagagtagg gtcccctcac ccaaaaaaaaa aaaaaaaaaa aaancttngg ggaaa 355
212 <210> SEQ ID NO: 14
213 <211> LENGTH: 658
214 <212> TYPE: DNA
215 <213> ORGANISM: Homo sapiens
217 <220> FEATURE:
218 <221> NAME/KEY: misc_feature
219 <222> LOCATION: 467, 527, 550, 579, 583, 600, 608, 616, 623, 625, 640, 655
220 <223> OTHER INFORMATION: n = A,T,C or G
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224 aatggacata ctgaagaaa gaatagacag tctagaagat caaattgaag aattctctaa 120
225 ggatacaatg caaatgacca aacagataat tagtaaagaa aggcaaagag atatatagga 180
226 gagatctaga agttgcaaca ttcgtttgat aggaattcca gaaaaggaga gttatgaga 240
227 tagggcagag gacataatta aagaaataat tgatgaaaac tttgcagaac taaagaaagg 300
228 ttcaagtctt gagattgtca gtgcttgctg agtacctagt aaaattgatg aaaagagact 360
229 gactcctaga cacatcttgg tgaaattttg gaattctagt gataaagaga aaataataag 420
230 ggcttctaga gagagaagag aaattaccta ccaaggaaca agaatcnggt tgacagcaga 480
231 cttatcctgg acacactgga tgctagaagt aaatgggagc caatgtnttc aaaggtcttg 540

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W--> 232 cttggaaaan ggcttttaaat cctagaatct tttttccanc canattgcct tttgatttan 600
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 236 <211> LENGTH: 713
 237 <212> TYPE: DNA
 238 <213> ORGANISM: Homo sapiens
 240 <220> FEATURE:
 241 <221> NAME/KEY: misc_feature
 242 <222> LOCATION: 505, 521, 546, 560, 563, 575, 594, 596, 626, 639, 689, 691
 243 <223> OTHER INFORMATION: n = A,T,C or G
 245 <400> SEQUENCE: 15
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 247 gacttacaaa tgggccaaag acaacttcaac ctcaaaaacca aagagaaatc tctgcttgca 120
 248 gagatacaaa gaaagtaact ctccctotta tgaaaagcaa ccaggaactc tactccagtt 180
 249 atgagggccca ctgatggtgt gggagagcta tcaagaagat tcttctaga cgtggtgcaa 240
 250 agacagtgag aaccacaggaa atcacattca tgggacactt gctcttaccg tcatcaccct 300
 251 ctattctatc tcaacttttg ccccatcaaa tctaatagata aacaaaagaa ggtaattaca 360
 252 tgtagaaaat caaagtgaat ggggaatgtg tgggtgtgaac ataaaagaag aaattgaaaa 420
 253 caatcaaaag tttctcagtg ctgctttccc gcactgtcat agaaatctct gatccaattc 480
 W--> 254 ttcatatgtc taacttccaa ggaancgggc taacagcaca nacataggat ccaaggcatt 540
 W--> 255 cttggngcgg aaaattaagn gngnggcccc ctttngaatt gggactgcaa ccahanggct 600
 W--> 256 gggaaatcct tatactcccc tcccgncggg ggctaattnc cattggtgaa acaatggttg 660
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 260 <211> LENGTH: 616
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 265 <221> NAME/KEY: misc_feature
 266 <222> LOCATION: 539, 551, 560, 563, 593, 601
 267 <223> OTHER INFORMATION: n = A,T,C or G
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 271 tacataaaac tagcagcaaa aagtatctag aaatctgtcg tgtgcaaata gttttcttcc 120
 272 caactatcat tcccatgggc ccaaataaat tttagaatct agtcccatcc ctttctaga 180
 273 caagctgctg tcaacaatct ccaagagaca aagtaagatt ggaagttaa ggacacgcac 240
 274 acaagacata tatataaaat tctctgaatg tgcaataaaa gaagtacttt gtaaaaagtt 300
 275 atgggcaaaa tgtacaaggg cctaaacctt gactaattga aatagcacca taacaaatga 360
 276 cctcaatact gtcaagtga cctacttaat aaaagtttta gaacaaggca caatacactt 420
 277 gaaaatctat tgcacttttag gaaatttttg ccgtcttct atgccactgt aaaaagatgg 480
 W--> 278 agcgttttga taccgcatt ctggacctcg ggccgcgacc caccgtaagg gcgaaattnc 540
 W--> 279 agcaccctt ngcgggcggn ttcttagtgg gatcccaact cggtaaccaa acnttggcgt 600
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 284 <212> TYPE: DNA
 285 <213> ORGANISM: Homo sapiens
 287 <220> FEATURE:
 288 <221> NAME/KEY: misc_feature

Use of n and/or Xaa has been detected in the Sequence Listing.
 Review the Sequence Listing to insure a corresponding
 explanation is presented in the <220> to <223> fields of
 each sequence using n or Xaa.

Use of n and/or Xaa has been detected in the Sequence Listing.
 Review the Sequence Listing to insure a corresponding
 explanation is presented in the <220> to <223> fields of
 each sequence using n or Xaa.

VERIFICATION SUMMARY

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Input Set : A:\493c1.app.txt

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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:42 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2

L:50 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2

L:110 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6

L:210 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13

L:230 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14

L:231 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14

L:232 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14

L:233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14

L:254 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15

L:255 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15

L:256 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15

L:257 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15

L:278 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16

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L:364 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21

L:377 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22

L:424 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25

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L:596 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35

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L:686 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41

L:687 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41